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Biochem J. 2003 Apr 1;371(Pt 1):39-48.

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PMID: 12513693 [PubMed - indexed for MEDLINE]

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     PCT Int. Appl., 124 pp.
SO
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     WO 2004039407
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JP 2004248668 A2 20040909 JP 2003-369723

20031030

PRAI JP 2002-320075 A 20021101

JP 2003-17892 A 20030127

AB It is intended to provide an apoptosis inducer or the like containing a compound

or its salt inhibiting the activity of a protein having an amino acid,

which is the same or substantially the same as the amino acid sequence of

HTRA3, its peptide fragment or a salt thereof or the expression of the

gene thereof.

- L3 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2005 ACS on STN
- AN 2004:927825 CAPLUS
- DN 142:293408
- TI Human HtrA proteases
- AU Dingwall, Colin; Holbrook, Joanna D.
- CS Alzheimer's Disease Research Department, Neurology & GI CEDD, GlaxoSmithKline, Harlow, CM19 5AW, UK
- SO Handbook of Proteolytic Enzymes (2nd Edition) (2004), Volume 2, 1476-1480.

Editor(s): Barrett, Alan J.; Rawlings, Neil D.; Woessner, J.
Fred.

Publisher: Elsevier, London, UK.

CODEN: 69GAQF; ISBN: 0-12-079610-4

DT Conference; General Review

LA English

AB A review. The human HtrA serine proteases (HtrA1-HtrA4) show extensive

homol. to the Escherichia coli HtrA (high-temperature requirement) protease,

also known as DegP, which is active in the periplasm of the bacterium and

is essential for bacterial tolerance of thermal, osmotic and oxidative

stress. The bacterial protein has the interesting property of acting as a

mol. chaperone at reduced temperature but acting as a protease at elevated

temps. The history, activity, specificity, protein structure, structural

chemical, preparation, and biol. aspects of HtrA proteases are briefly discussed.

RE.CNT 21 THERE ARE 21 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

L3 ANSWER 3 OF 3 MEDLINE on STN

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DN PubMed ID: 12513693

TI Identification and cloning of two isoforms of human

high-temperature

requirement factor A3 (HtrA3), characterization of its genomic structure

and comparison of its tissue distribution with HtrA1 and HtrA2.

AU Nie Gui-Ying; Hampton Anne; Li Ying; Findlay Jock K; Salamonsen Lois A

CS Prince Henry's Institute of Medical Research, P.O. Box 5152, 246 Clayton

Road, Clayton, Victoria 3168, Australia...

guiying.nie@med.monash.edu.au

SO Biochemical journal, (2003 Apr 1) 371 (Pt 1) 39-48.

Journal code: 2984726R. ISSN: 0264-6021.

CY England: United Kingdom

DT Journal; Article; (JOURNAL ARTICLE)

LA English

FS Priority Journals

EM 200305

ED Entered STN: 20030321

Last Updated on STN: 20030523

Entered Medline: 20030522

AB In the present study, we identified an additional member of the human

high-temperature requirement factor A (HtrA) protein family, called

pregnancy-related serine protease or HtrA3, which was most highly expressed in the heart and placenta. We cloned the full-length sequences

of two forms (long and short) of human HtrA3 mRNA,

located the gene on chromosome 4p16.1, determined its genomic structure

and revealed how the two mRNA variants are produced through alternative

splicing. The alternative splicing was also verified by Northern blotting. Four distinct domains were found for the long form HtrA3

protein: (i) an insulin/insulin-like growth factor binding
domain, (ii) a

Kazal-type S protease-inhibitor domain, (iii) a trypsin protease domain

and (iv) a PDZ domain. The short form is identical to the long form

except it lacks the PDZ domain. Comparison of all members of human HtrA

proteins, including their isoforms, suggests that both isoforms of HtrA3

represent active serine proteases, that they may have different substrate

specificities and that HtrA3 may have similar functions to HtrA1. All

three HtrA family members showed very different mRNA-expression patterns

in 76 human tissues, indicating a specific function for each.

Interestingly, both HtrA1 and HtrA3 are highly expressed in the placenta.

Identification of the tissue-specific function of each HtrA family member

is clearly of importance.

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